

FIG. 1

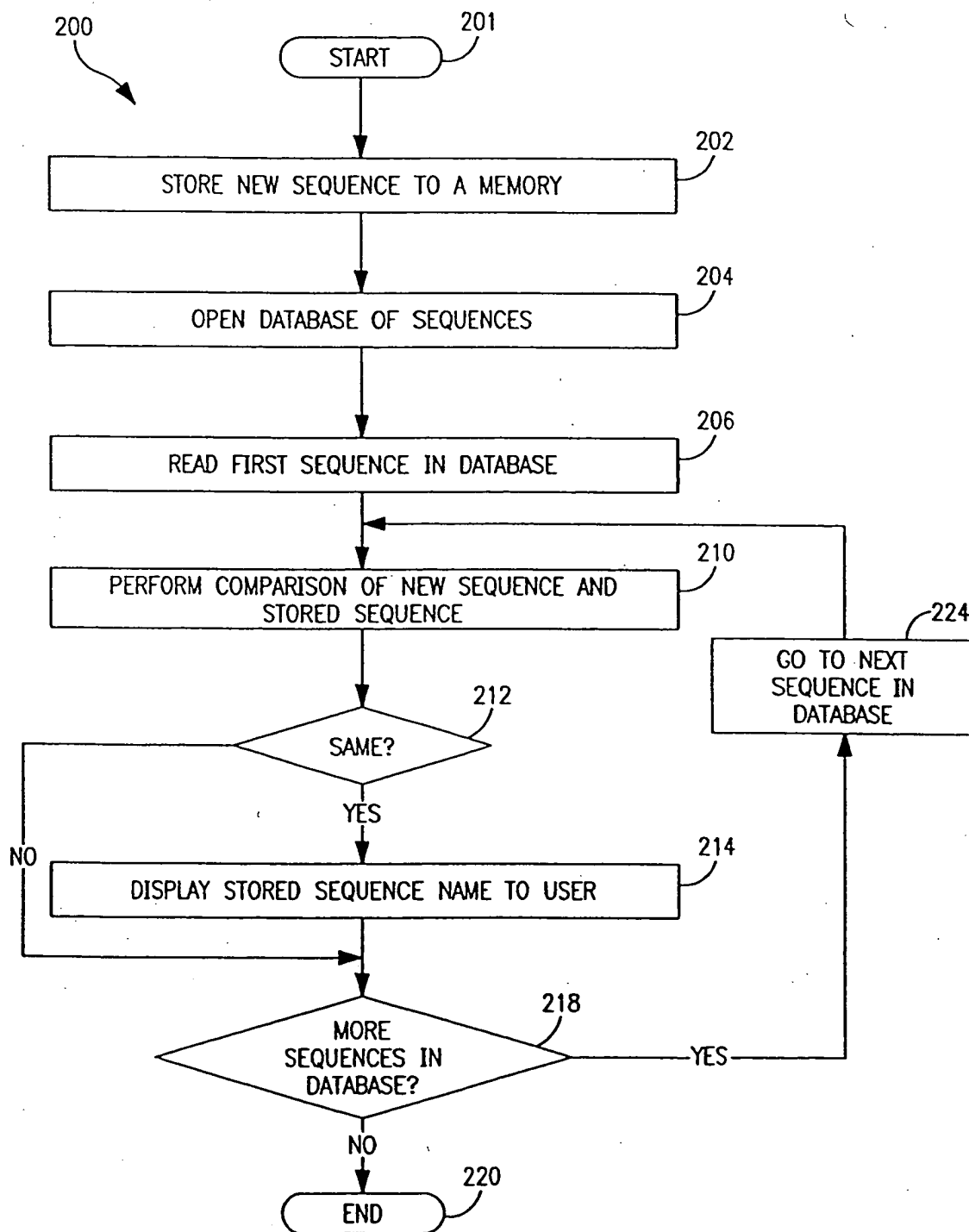


FIG. 2

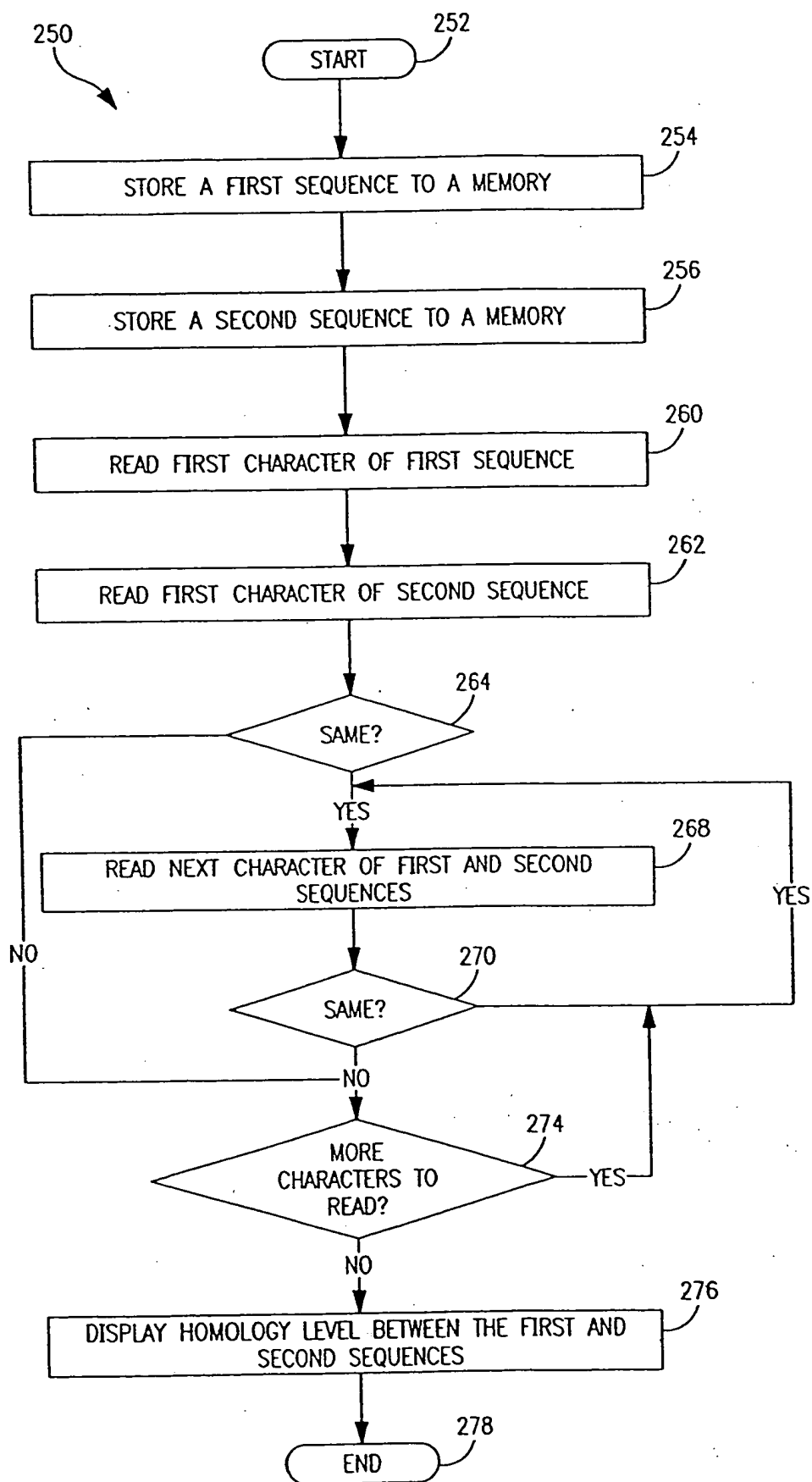


FIG. 3

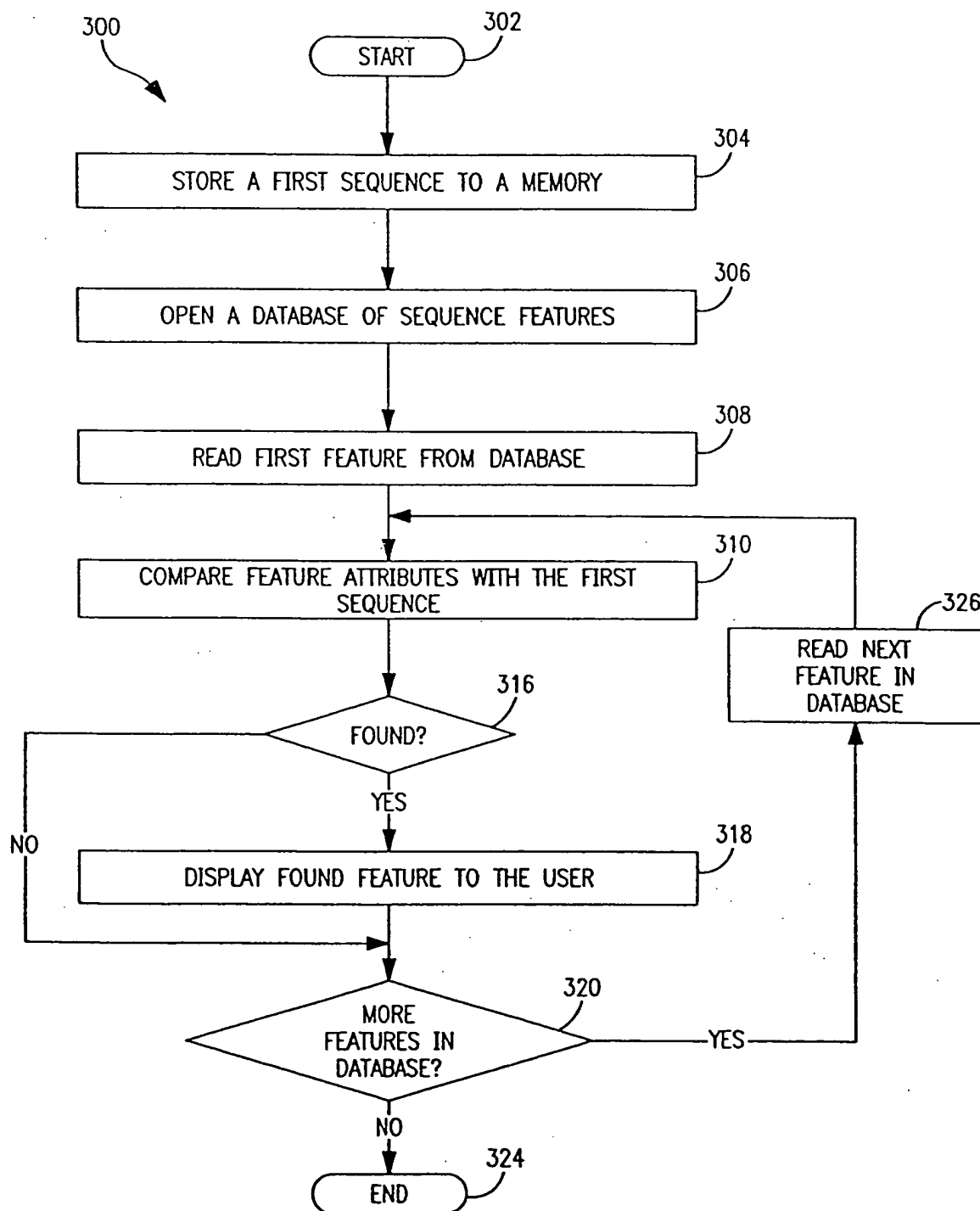


FIG. 4



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**Thermococcus GU5L5 Amidase**

1 ATG ACC GGC ATC GAA TGG AAC CAC GAG ACC TTT TCT AAG TTC GCC TAC CTG GGC GAC CCG 60  
1 Met Thr Gly Ile Glu Trp Asn His Glu Thr Phe Ser Lys Phe Ala Tyr Leu Gly Asp Pro 20  
  
61 AGG ATA CGG GGA AAC TTA ATC GCG TAC ACC CTG ACG AAG GCC AAC ATG AAG GAC AAC AAG  
120  
21 Arg Ile Arg Gly Asn Leu Ile Ala Tyr Thr Leu Thr Lys Ala Asn Met Lys Asp Asn Lys 40  
  
121 TAC GAG AGC ACG GTT GTT GAA GAC CTT GAA ACG GGC TCA AGG CGC TTC ATC GAG AAC 180  
41 Tyr Glu Ser Thr Val Val Glu Asp Leu Glu Thr Gly Ser Arg Phe Ile Glu Asn 60  
  
181 GCC TCA ATG CCG AGG ATT TCG CCA GAC GGC AGA AAG CTC GCC TTC ACC TGC TTT AAC GAG 240  
61 Ala Ser Met Pro Arg Ile Ser Pro Asp Gly Arg Lys Leu Ala Phe Thr Cys Phe Asn Glu 80  
  
241 GAG AAG AAG GAG ACC GAG ATA TGG GTG GCC GAT ATC CAG ACC CTG AGC GCC AAG AAA GTC  
300  
81 Glu Lys Lys Glu Thr Glu Ile Trp Val Ala Asp Ile Gln Thr Leu Ser Ala Lys Lys Val 100

FIGURE 5 A



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301 CTC TCA ACT AAA AAC GTC CGC TCG ATG CAG TGG AAC GAC GAT TCA AGG AGA CTC TTA GTT 360  
101 Leu Ser Thr Lys Asn Val Arg Ser Met Gln Trp Asn Asp Asp Ser Arg Arg Leu Val 120

361 GTC GGC TTC AAG AGG AGG GAC GAT GAG GAC TTC GTC TTT GAC GAC GAC GTC CCG GTC TGG 420  
121 Val Gly Phe Lys Arg Arg Asp Asp Glu Asp Phe Val Phe Asp Asp Val Pro Val Trp 140

421 TTC GAC AAT ATG GGA TTC TTT GAT GGA GAG AAG ACG ACG TTC TGG GTT CTT GAC ACT GAG 480  
141 Phe Asp Asn Met Gly Phe Phe Asp Gly Glu Lys Thr Thr Trp Val Leu Asp Thr Glu 160

481 GCC GAG GAG ATA ATC GAG CAG TTC GAG AAG CCG AGG TTT TCG AGT GGC CTC TGG CAC GGC 540  
161 Ala Glu Glu Ile Ile Glu Gln Phe Glu Lys Pro Arg Phe Ser Ser Gly Leu Trp His Gly 180

541 GAT GCG ATA GTT GTG AAC GTC CGC CAC CGC GAG GGG AGC AAG CCT GCC CTG TTC AAG TTC 600  
181 Asp Ala Ile Val Val Asn Val Pro His Arg Glu Gly Ser Lys Pro Ala Leu Phe Lys Phe 200

601 TAC GAC ATA GTC CTA TGG AAG GAC GCG GAG GAA GAG AAG CTC TTC GAG AGG GTC TCC TTC 660  
201 Tyr Asp Ile Val Leu Trp Lys Asp Gly Glu Glu Lys Leu Phe Glu Arg Val Ser Phe 220

661 GAG GCG GTT GAC TCC GAC GGA AAG AGA ATA CTC CTG AGG GGC AAG AAA AAA AAG CGG TTC 720

721 Glu Ala Val Asp Ser Asp Gly Lys Arg Ile Leu Leu Arg Gly Lys Lys Lys Arg Phe 240

FIGURE 5 B



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721 ATC AGC GAG CAC GAC TGG CTG TAC CTC TGG GAC GGC GAG CTT AAA CCG ATC TAC GAG GGC 780  
241 Ile Ser Glu His Asp Trp Leu Tyr Leu Trp Asp Gly Glu Leu Lys Pro Ile Tyr Glu Gly 260

781 CCG CTC GAC GTC TGG GAA GCC AAG CTC ACG GAA GGA AAG GTC TAC TTC CTC ACT CCA GAT 840  
261 Pro Leu Asp Val Trp Glu Ala Lys Leu Thr Glu Gly Lys Val Tyr Phe Leu Thr Pro Asp 280

841 GCG GGC AGG GTA AAC CTC TGG CTC TGG GAC GGG AAG GCC GAG CGT GTT ACC GGC GAC 900  
281 Ala Gly Arg Val Asn Leu Trp Leu Trp Asp Gly Lys Ala Glu Arg Val Val Thr Gly Asp 300

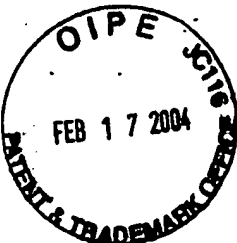
901 CAC TGG ATT TAC GGG CTT GAC GTC AGC GAT GGC AAA GCA TTG CTC CTC ATC ATG ACC GCC 960  
301 His Trp Ile Tyr Gly Leu Asp Val Ser Asp Gly Lys Ala Leu Leu Ile Met Thr Ala 320

961 ACG AGG ATA GGC GAG CTC TAC CTC TAC GAC GGC GAG CTG AAA CAG GTC ACC GAA TAC AAC  
1020  
321 Thr Arg Ile Gly Glu Leu Tyr Leu Tyr Asp Gly Glu Leu Lys Gln Val Thr Glu Tyr Asn 340

1021 GGG CCG ATA TTC AGG AAG CTC AAG ACC TTC GAG CCG AGG CAC TTC CGC TTC AAG AGC AAA  
1080  
341 Gly Pro Ile Phe Arg Lys Leu Lys Thr Phe Glu Pro Arg His Phe Arg Phe Lys Ser Lys 360

1081 GAC CTC GAG ATA GAC GGC TGG TAC CTC ACG CCG GAG GTT AAA GAG GAG AAG GCC CCG GTG  
1140  
361 Asp Leu Glu Ile Asp Gly Trp Tyr Leu Arg Pro Glu Val Lys Glu Glu Lys Ala Pro Val 380

FIGURE 5c



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1141 ATA GTC TTC GTC CAC GGC GGG CCG AAG GGC ATG TAC GGA CAC CGC TTC GTC TAC GAG ATG  
1200  
381 Ile Val Phe Val His Gly Gly Pro Lys Gly Met Tyr Gly His Arg Phe Val Tyr Glu Met 400  
  
1201 CAG CTG ATG GCG AGC AAG GGC TAC TAC GTC GTC TTC GTG AAC CCG CGC GGC AGC GAC GGC  
1260  
401 Gln Leu Met Ala Ser Lys Gly Tyr Tyr Val Val Phe Val Asn Pro Arg Gly Ser Asp Gly 420  
  
1261 TAT AGC GAA GAC TTC GCG CTC CGC GTC CTG GAG AGG ACT GGC TTG GAG GAC TTT GAG GAC  
1320  
421 Tyr Ser Glu Asp Phe Ala Leu Arg Val Leu Glu Arg Thr Gly Leu Glu Asp Phe Glu Asp 440  
  
1321 ATA ATG AAC GGC ATC GAG GAG TTC TTC AAG CTC GAA CCG CAG GCC GAC AGG GAG CGC GTT  
1380  
441 Ile Met Asn Gly Ile Glu Glu Phe Phe Lys Leu Glu Pro Gln Ala Asp Arg Glu Arg Val 460  
  
1381 GGA ATA ACG GGC ATA AGC TAC GGC GGC TTC ATG ACC AAC TGG GCC TTG ACT CAG AGC GAC  
1440  
461 Gly Ile Thr Gly Ile Ser Tyr Gly Gly Phe Met Thr Asn Trp Ala Leu Thr Gln Ser Asp 480  
  
1441 CTC TTC AAG GCA GGA ATA AGC GAG AAC GGC ATA AGC TAC TGG CTC ACC AGC TAC GCC TTC  
1500  
481 Leu Phe Lys Ala Gly Ile Ser Glu Asn Gly Ile Ser Tyr Trp Leu Thr Ser Tyr Ala Phe 500

FIGURE 5 D





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1501 TCG GAC ATA GGG CTC TGG TAC GAC GTC GAG GTC ATC GGG CCA AAT CCG TTA GAG AAC GAG  
1560  
501 Ser Asp Ile Gly Leu Trp Tyr Asp Val Glu Val Ile Gly Pro Asn Pro Leu Glu Asn Glu 520  
1561 AAC TTC AGG AAG CTC AGC CCG CTG TTC TAC GCT CAG AAC GTG AAG GCG CCG ATA CTC CTA  
1620  
521 Asn Phe Arg Lys Leu Ser Pro Leu Phe Tyr Ala Gln Asn Val Lys Ala Pro Ile Leu Leu 540  
1621 ATC CAC TCG CTT GAG GAC TAC CGC TGT CCG CTC GAC CAG AGC CTT ATG TTC TAC AAC GTG  
1680  
541 Ile His Ser Leu Glu Asp Tyr Arg Cys Pro Leu Asp Gln Ser Leu Met Phe Tyr Asn Val 560  
1681 CTC AAG GAC ATG GGC AAG GAA GCC TAC ATA GCG ATA TTC AAG CGC GGC GCC CAC GGC CAC  
1740  
561 Leu Lys Asp Met Gly Lys Glu Ala Tyr Ile Ala Ile Phe Lys Arg Gly Ala His Gly His 580  
1741 AGC GTC CGC GGA AGC CCG AGG CAC AGG CCG AAG CGC TAC AGG CTC TTC ATA GAG TTC TTC  
1800  
581 Ser Val Arg Gly Ser Pro Arg His Arg Pro Lys Arg Tyr Arg Leu Phe Ile Glu Phe Phe 600  
1801 GAG CGC AAG CTC AAG AAG TAC GAG GAG GGC TTT GAG GTA GAG AAG ATA CTC AAG GGG AAT  
1860  
601 Glu Arg Lys Leu Lys Lys Tyr Glu Glu Gly Phe Glu Val Glu Lys Ile Leu Lys Gly Asn 620  
1861 GGG AAC TGA 1869  
621 Gly Asn End 623

FIGURE 5 F

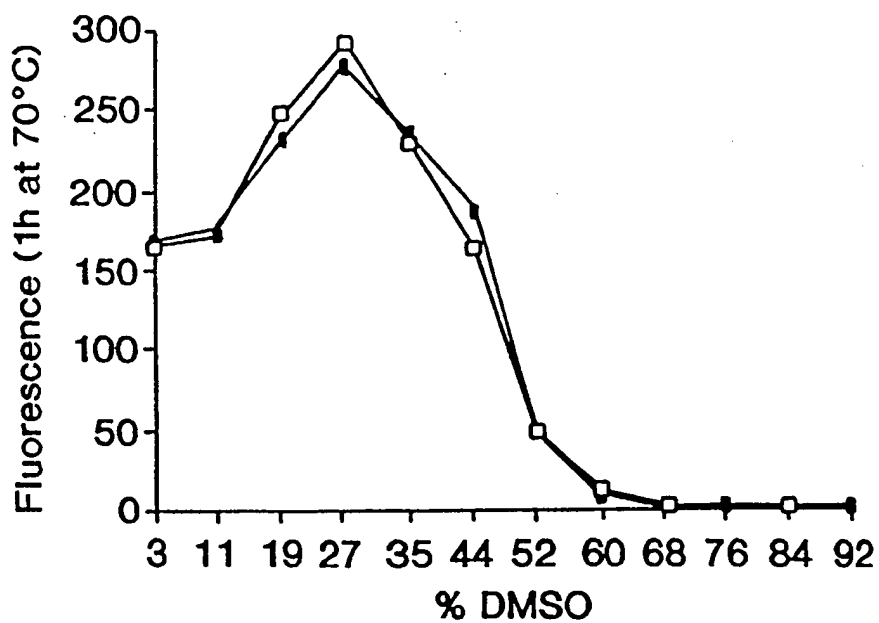


FIGURE 6

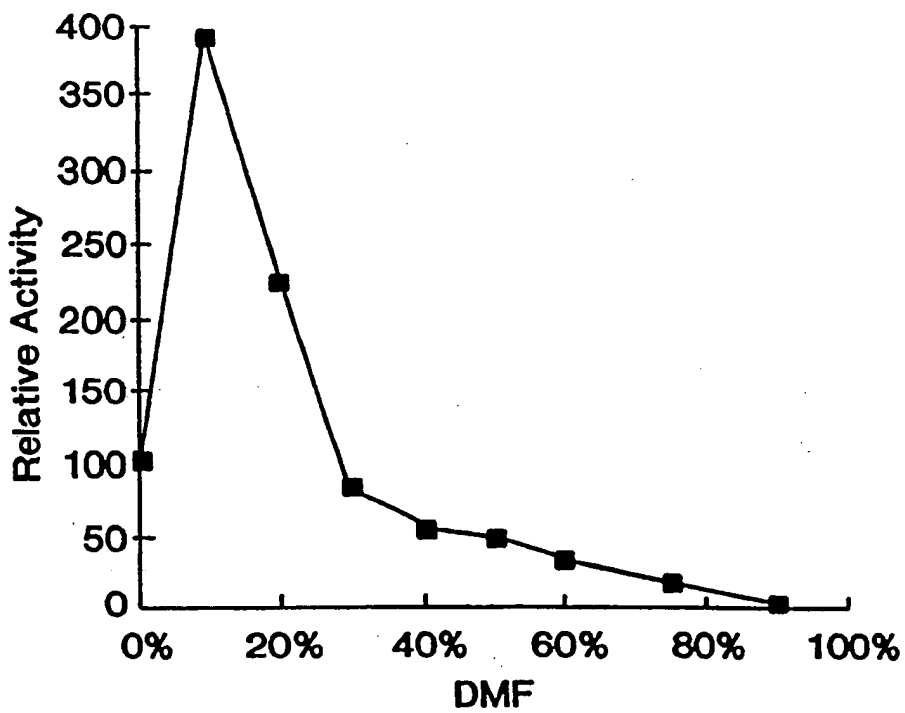


FIGURE 7